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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tissue Procurement. Jeffrey Medeiros, M.D., Michael R. Emmert-Buck, M.D., Ph.D. Emmert-Buck, M.D., Ph.D. CDNA Library Preparation: Stratagene, Inc. CDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ACCESSION: AA577530
ACCESSION: AA535711
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 431)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCI-CGAP http://www.ncbi.nlm.nih.gov/nciogap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             1: /home/sdavid/sdavid-tmp/mar03/yu602/aa577530.gb_est1:*
2: /home/sdavid/sdavid-tmp/mar03/yu602/aa535711.gb_est1:*
                                                                                           March 13, 2003, 13:29:58; Search time 2 Seconds (without alignments) 3.694 Million cell updates/sec
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Compugen Ltd.
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   GenCore version (c) 1993 - 2003
                                                                 sw model
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Listing first 2 summaries
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AA535711
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Gapop 10.0 , Gapext 0.5
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length: 2000000000
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EST 21-AUG-1997
Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
www.bio.llnl.gov/bbrp/image/image.html
Seq primer: -40ml3 fwd.ET from Amersham
High quality sequence stop: 386.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 478)
NCI-CAP http://www.ncbi.nlm.nih.gov/ncicgap.
NCI-CAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /issue_type="colon tumor"
/lab_host="SoLR (kanamycin resistant)"
/note="Organ: colon; Vector: Bluescript SK-; Site_l:
/site_l: Xhol; Cloned unidirectionally. Primer: Oli
dT. Pooled colon tumors. 5' adaptor sequence: 5'
GAATTCGGCACGAG 3' 3' adaptor sequence: 5'
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.nf88d08.s1 NCI_CGAP_CO3 Homo sapiens cDNA clone IMASE.
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0; Mismatches
                                                                                                                                                         /db_xref="taxon:9606"
/clone="IMAGE:1083839"
/clone_lib="NCI_CGAP_Col2"
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Pred. No. 0:
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AA535711.1 GI:2279964
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Best Local Similarity
Matches 408; Conserv
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    48
ACAATGTTTTATGGCCTAAATTTATAATAAATAAATGCACAAAACTTTG 2467
                                                                                                                                                                                 TTAAAAAGGAGAAGATAACTGAGGATTTTAAAAAGAAGCCATGGAAAAACTTCCTAGTAA 2359
                                                                                                                                                                                                                                                                                                                                                                                                             AAGGTCACATATCCTGGAGCAGGAAAACTTCCACACATCATTGGAGGATCAGATCTAATA 289
                                                                                                                         GCATCTACTTCAGGCCAACAAGGTTATATGAATATATAGTGTATAGAAGCGATTTAAGTT 2419
                                                                                                                                                            Tumor Gene Index
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
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Insert Length: 2473 Std Error: 0.00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cDNA Library Preparation: M. Bento Soares, Ph.D. cDNA Library Arraying: Greg Lennon, Ph.D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone="IMAGE:926991"
/clone_lib="NCI_CGAP_Co3"
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db_xref="taxon:9606"
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